SEQUENCE ID LISTING

SEQUENCE ID NUMBER 1

SEQ. ID NO.: 1 is a transducisome protein (fly) amino acid sequence.

MVQFLGKQGT AGELIHMVTL DKTGKKSFGI CIVRGEVKDS

- 5 PNTKTTGIFI
 KGIVPDSPAH LCGRLKVGDR ILSLNGKDVR NSTEQAVIDL
 IKEADFKIEL
 EIQTFDKSDE QQAKSDPRSN GYMQAKNKFN QEQTTNNNAS
 GGQGMGQGQG
- 10 QGQGMAGMNR QQSMQKRNTT FTASMRQKHS NYADEDDEDT RDMTGRIRTE
 AGYEIDRASA GNCKLNKQEK DRDREQEDEF GYTMAKINKR YNMMKDLRRI
 EVQRDASKPL GLALAGHKDR QKMACFVAGV DPNGALGSVD
- 15 IKPGDEIVEV
 NGNVLKNRCH LNASAVFKNV DGDKLVMITS RRKPNDEGMC
 VKPIKKFPTA
 SDETKFIFDQ FPKARTVQVR KEGFLGIMVI YGKHAEVGSG
 IFISDLREGS
- 20 NAELAGVKVG DMLLAVNQDV TLESNYDDAT GLLKRAEGVV
 TMILLTLKSE
 EAIKAEKAAE EKKKEEAKKE EEKPQEPATA EIKPNKKILI
 ELKVEKKPMG
 CHRLRRQKQP CHDWLCNHPR LSGGQVAADK RLKIFDHICD
- 25 INGTPIHVGS

 MTTLKVHQLF HTTYEKAVTL TVFRADPPEL EKFNVDLMKK

 AGKELGLSLS

 PNEIGCTIAD LIQGQYPEID SKLQRGDIIT KFNGDALEGL

 PFQVCYALFK

GANGKVSMEV TRPKPTLRTE APKA

SEQUENCE ID NUMBER 2

SEQ. ID NO. 2 is a transducisome protein (fly) nucleotide sequence.

(InaD) mRNA, complete cds InaD cDNA

- 5 ATGGTTCAGTTCCTGGGCAAACAGGGCACCGCGGGTGAGCTCATTCA
 CATGGTGACCCTGGACAAG
 ACGGGCAAGAAGTCCTTCGGCATCTGCATAGTGCGCGGCGAGGTGAA
 GGATTCGCCCAACACCAAGACAA
 CCGGCATCTTCATCAAGGGCATTGTGCCCGACAGTCCCGCGCATCTGT
- 10 GTGGTCGCCTAAAGGTTGGCGA
 TCGGATCCTCTCGCTCAACGGAAAGGATGTGCGCAACTCCACCGAAC
 AGGCGGTCATCGATCTCATCAAG
 GAGGCGGACTTCAAGATCGAGCTTGAGATTCAGACCTTCGACAAGAG
 CGATGAGCAGCAGGCCAAGTCAG
- 15 ATCCGCGGAGCAATGGCTACATGCAGGCCAAGAACAAGTTCAATCAG
 GAGCAGACCACCAACAACAATGC
 GTCCGGAGGTCAGGGAATGGGCCAAGGTCAGGGTCAGGGA
 ATGGCTGGCATGAACCGGCAGCAA
 TCGATGCAGAAGCGGAATACCAGATTCACGGCCTCGATGCGTCAGAA
- 20 GCATAGTAACTACGCCGACGAGG
 ATGACGAGGACACCCGGGACATGACCGGTCGCATTCGCACGGAGGCG
 GGTTATGAGATCGATCGAGCCTC
 CGCCGGTAATTGCAAACTTAATAAGCAGGAAAAGGATCGCGACAAG
 GAGCAGGAAGATGAATTTGGCTAC
- 25 ACGATGGCTAAGATCAACAAGCGGTACAACATGATGAAGGATCTGCG
 CAGGATCGAGGTCCAGAGGGACG
 CCAGCAAGCCACTGGGACTCGCACTCGCTGGCCACAAGGACCGCCAG
 AAGATGGCCTGCTTTGTTGCCGG
 TGTGGATCCCAACGGAGCATTGGGCAGCGTGGACATTAAGCCGGGCG
- 30 ACGAGATCGTCGAGGTCAACGGC

 AATGTGCTTAAGAATCGCTGCCACTTGAACGCCTCCGCCGTGTTCAAG

 AGCGTGGATGGGGATAAGCTCG

 TGATGATCACCTCGCGACGCAAGCCCAACGATGAGGGCATGTGCGTC

 AAGCCCATCAAAAAGTTCCCCAC
- 35 CGCGTCTGATGAGACTAAGTTTATCTTCGACCAGTTTCCCAAGGCGCG

CACGGTGCAGGTGCGCAAGGAG GGTTTCCTGGGCATCATGGTCATCTATGGCAAGCACGCTGAGGTGGG CAGTGGCATTTTCATCTCGGATC TGAGAGAGGGATCGAATGCCGAGTTGGCGGGCGTGAAAGTGGGCGA 5 CATGCTGCTGGCCGTTAATCAGGA TGTAACACTGGAATCCAACTACGATGATGCTACTGGACTGCTTAAAC GTGCCGAGGGCGTAGTGACCATG ATTCTATTGACTCTCAAGAGCGAGGGGGGGGATAAAGGCTGAGAAGGC AGCGGAAGAGAAAAAGAAGGAG& 10 AGGCCAAGAAAGAGGAGGAAAA&CCACAGGAACCCGCCACAGCCGA GATCAAGCCGAACAAAAAGATACT CATTGAGTTGAAGGTGGAAAAGAAGCCAATGGGCGTCATCGTCTGCG GCGGCAAGAACAACCATGTCA#G ACTGGCTGTGTAATCACCCAC#TFTATCCGGAGGGACAAGTGGCAGC 15 CGACAAGCGCCTCAAGATCTTTG ACCACATTTGCGATATAAATGGTACGCCAATCCACGTGGGATCCATG ACGACACTGAAGGTCCATCAGTT ATTCCACACCACATACGAGAAG&CGGTCACCCTAACGGTCTTCCGCG CTGATCCTCCGGAACTGGAAAAG 20 TTTAACGTTGACCTTATGAAAAAAGCAGGCAAGGAGCTGGGCCTGTC GCTGTCTCCCAACGAAATTGGAT GCACCATCGCGGACTTGATTCAAGGACAATACCCGGAGATTGACAGC AAACTGCAGCGCGGCGATAT TAT CACCAAATTCAATGGCGATGCCTTGGAGGGTCTTCCGTTCCAGGTGTG 25 CTACGCCTTGTTCAAGGGAGCC AACGGCAAGGTATCGATG\$AAGTGACACGACCCAAGCCCACTCTACG

TACGGAGGCACCCAAGGCCTAGA

GACGATCCTCATTCTCCTCTCCGTAGCGAAGCAGT